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RAW SEQUENCE LISTING DATE: 04/08/2002 PATENT APPLICATION: US/10/002,292A TIME: 15:15:39

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\04082002\J002292A.raw

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      3 -:110> AFPLICANT: Ward, Brian
             Snyder, Leanne
      5
             Li, Chuan
             Song, Keming
             Opper, Kristen
             Uder, Stephanie
             Hernan, Ron
     11 <120> TITLE OF INVENTION: RECOMBINANT DNA PROCESSES USING A dNTP MIXTURE CONTAINING
MODIFIED
    12
             NUCLEOTIDES
     14 <130> FILE REFERENCE: SGM 6938.1
    16 <140> CURRENT APPLICATION NUMBER: US 10/002,292A
C--> 17 <141> CURRENT FILING DATE: 2002-03-25
    19 <150> PRIOR APPLICATION NUMBER: US 60/325,612
    20 <151 > PRIOR FILING DATE: 2001-09-28
    22 < 160 > NUMBER OF SEQ ID NOS: 14
    24 <170 > SOFTWARE: PatentIn version 3.1
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    29 - 213> ORGANISM: Lambda phage
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    32 <221> NAME/KEY: misc_feature
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F4 - 223 - OTHER INFORMATION: Primer for Polymerase Chain Reaction (PCR) €7 +400 + SEQUENCE: 3 An invariguate egatgagite gigiteegiae aactg 35 71 + 110 + SEQ ID NO. 4 L + 211 + LENGTH: 35 73 - 212 - TYPE: DNA 74 - 213 - ORGANISM Lambda phage The + 220 + FEATURE: 77 + L21 + NAME/KEY = misc_feature 78 < 222 > LOCATION: (1)..(35) 79 < 223 > OTHER INFORMATION: Primer for Polymerase Chain Reaction (PCR) 82 <400> SEQUENCE: 4 83 gcacqtotag aggttatoga aatcagocac agogo 35 86 <210 > SEQ ID NO: 5 87 <211> LENGTH: 23 88 <212> TYPE: DNA 89 <213> ORGANISM: Lambda phage 91 <220> FEATURE: 92 <221> NAME/KEY: misc_feature 93 <222> LOCATION: (1)..(23) 94 <223 > OTHER INFORMATION: Primer for Polymerase Chain Reaction (PCR) 97 <400> SEQUENCE: 5 98 caqteacgae gttgtaaaac gae 23 101 <210> SEQ ID NO: 6 102 <211> LENGTH: 23 103 <212> TYPE: DNA 104 <213> ORGANISM: Lambda phage 106 <220> FEATURE: 107 <221> NAME/KEY: misc_feature 108 <222> LOCATION: (1)..(23) 109 <223> OTHER INFORMATION: Primer for Polymerase Chain Reaction (PCR) 112 <400> SEQUENCE: 6 113 cacaggaaac agctatgacc atg 23 116 <210> SEQ ID NO: 7 117 <211> LENGTH: 23 118 - 212> TYPE: DNA 119 <213> ORGANISM: Lambda phage 121 < 220 > FEATURE: 122 <221> NAME/KEY: misc_feature 123 <222> LOCATION: (1)..(23) 124 <223> OTHER INFORMATION: Primer for Polymerase Chain Reaction (PCR) 127 <400> SEQUENCE: 7 128 ggcaatcagt toatottteg tea 23 131 <210> SEQ ID NO: 8 132 <211> LENGTH: 22 133 <212> TYPE: DNA 134 - 213 > ORGANISM: Lambda phage 136 <220> FEATURE: 137 <221> NAME/KEY misc_feature

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     141 KL11 + LENGTH: 558
     148 KILDE TYPE: DNA
     149 KUlta ORGANISM: Escherichia coli
     151 -:400% SEQUENCE: 9
     152 agettotega gatgeetgtt etggaaaaee gggetgetea gggegatatt aetgeaeeeg
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     154 graytgeteg cegittaacg ggigateaga eigeogetet gegigattet eitagegata
                                                                                120
     15% aacctgcaaa aaatattatt ttgctgattg gcgatgggat gggggactcg gaaattactg
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     15% occcacqtaa ttatgeegaa ggtgegggeg getttttaa aggtatagat geettacege
     160 thaccoggos atacacteae tatgegotga ataaaaaaac eggeaaaceg gactaegtea
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     162 conactogge typeateagea accepting caaceggity caaaacetat aacggegege
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     164 tagagegtega tatteaegaa aaagateaee caaegattet ggaaatggea aaageegeag
     166 qtetggegae eggtaaegtt tetaeegeag agttgeagga tgeeaegeee getgegetgg
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     174 - 223> OTHER INFORMATION: TAG USED TO PURIFY RECOMINANT PROTEIN
     181 - 220> FEATURE:
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     183 + 2225 LOCATION: (1)..(8)
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     193 - 2105 SEQ ID NO: 11
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     200
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     203 <221> NAME/KEY: promoter
     204 < 222 > Location: (1)...(60)
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     206
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     214 -..11> LENGTH: 60
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- 216 <213 ORGANISM: Artificial Sequence
- 218 122(→ FEATURE:
- $219{ ext{ o}}223{ ext{ o}}$ OTHER INFORMATION: complimentary sequence to SEQ ID 11 containing cohesive end

to en

- hance directional ligation to a gene for amplification 220
- 222 <120% FEATURE:
- 223 <221 × NAME/KEY: promoter
- 224 R222 F LOCATION: (1)..(60)
- 225 223 OTHER INFORMATION: complimentary sequence to SEQ ID 11 containing cohesive end

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- hance directional ligation to a gene for amplification 226
- 129 <4008 SEQUENCE: 12
- 230 octaogatta tgctgagtga tatccctctt cccggtggta cctgatgttt ctgctactgc 60
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- 235 <212> TYPE: DNA
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- 242 <220> FEATURE:
- 243 <221> NAME/KEY: primer_bind
- 244 <222> LOCATION: (1)..(50)
- 245 < 223 > OTHER INFORMATION: complimentary sequence to SEQ ID 14 containing a cohesive end to
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 - 249 <400> SEQUENCE: 13
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 - 255 <212> TYPE: DNA
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- 263 <221> NAME/KEY: primer_bind
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- ese sequences into PCR amplification products 266
- 269 <400> SEQUENCE: 14
- 270 aacttttgat tggtatgcag tacacgggtg gtcggaacag gattat

50

VERIFICATION SUMMARY

DATE: 04/08/2002 PATENT APPLICATION: US/10/002,292A TIME: 15:15:40

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